

a) at least [120] 394 contiguous nucleotides of the open reading frame of SEQ ID NO: 1; or

b) at least 17 contiguous nucleotides of the open reading frame of SEQ ID NO: 3;

to form a duplex, wherein detection of said duplex indicates the presence of said polynucleotide.

9. (Three times amended) A kit for the detection of a polynucleotide, comprising a compartment containing a probe that hybridizes, under stringent hybridization wash conditions of at least 65°C, less than about 150 mM salt , to:

a) at least [120] 394 contiguous nucleotides of the open reading frame of SEQ ID NO: 1; or

b) at least 17 contiguous nucleotides of the open reading frame of SEQ ID NO: 3;

to form a duplex, wherein detection of said duplex indicates the presence of said polynucleotide.

21. (Twice amended) The method of claim 8 wherein said probe hybridizes to at least [120] 394 contiguous nucleotides of said open reading frame of SEQ ID NO: 1.

22. (Canceled) The method of claim 21 wherein said probe hybridizes to at least 140, 175, 200, or 300 contiguous nucleotides of said open reading frame of SEQ ID NO: 1.

25. (Twice amended) The kit of claim 9 wherein said probe hybridizes to at least [120] 394 contiguous nucleotides of said open reading frame of SEQ ID NO: 1.

26. (Canceled) The kit of claim 25 wherein said probe hybridizes to at least 140, 175, 200, or 300 contiguous nucleotides of said open reading frame of SEQ ID NO: 1.